

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re application of

Paulo César PEREGRINO FERREIRA et al.

Serial No. 09/759,281

BOX PCT

Filed January 16, 2001

Attention: DO/EO

METHOD AND COMPOSITION FOR THE DIAGNOSIS OF EQUINE INFECTIOUS ANEMIA VIRUS DISEASE BY USING THE RECOMBINANT CAPSID PROTEIN

## SUBMISSION UNDER 37 C.F.R. §821 FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Commissioner for Patents

Washington, D.C. 20231

Sir:

Applicants attach to the present paper a "Sequence Listing", as required by Rule 821(c), and a copy of the "Sequence Listing" in computer readable form, as required by Rule 821(e).

Applicants hereby state that the content of the attached paper and computer readable copy of the Sequence Listing are the same.

Respectfully submitted,

YOUNG & THOMPSON

Andrew J. Patch

Attorney for Applicants Registration No. 32,925 745 South 23rd Street Arlington, VA 22202 Telephone: 521-2297

June 6, 2001 Ref. 41826 CIP



SEQUENCE LISTING

**GENERAL INFORMATION:** 

(i)

APPLICANT: PEREGRINO FERREIRA, Paulo;

5 GESSIEN KROON, Erna;

PIMENTA DOS REIS, Karlisson Jennner;

BIAS FORTES FERRAZ, Isabella;

CERQUEIRA LEITE, Romulo.

(ii)

TITLE OF INVENTION: Method and composition for the diagnosis of equine infectious anemia virus disease by using the recombinant capsid protein virus (p26)

(iii)

NUMBER OF SEQUENCES: 1

15 (iv)

**CORRESPONDENCE ADDRESS:** 

(A)

ADDRESSEE: Universidade Federal de Minas Gerais - CTIT

(B)

20 STREET: Avenida Antônio Carlos, 6627 Bairro São Francisco

(C)

CITY: Belo Horizonte

(D)

STATE: Minas Gerais

25 (E)

COUNTRY: BRAZIL

(F)

ZIP: 31270-901

(v)

30 COMPUTER READABLE FORM:

(A)

MEDIUM TYPE: diskette - 3.50 inch, 1.44 Mb storage

(B)

COMPUTER: IBM compatible

(C)

5 OPERATING SYSTEM: Windows 98

(D)

SOFTWARE: Office premium

(vi)

**CURRENT APPLICATION DATA:** 

10 (A)

APPLICATION NUMBER: U.S. 09/331.262

(B)

FILING DATE:

(C)

15 CLASSIFICATION: C12Q1/70

(vii)

PRIOR APPLICATION DATA

(A)

APPLICATION NUMBER: PI 9606273-8

20 **(B)** 

FILING DATE: 18-DEC-1996

(2)

**INFORMATION FOR SEQ ID N0:1:** 

(i)

25 SEQUENCE CHARACTERISTICS:

(A)

LENGHT: 252 amino acids

(B)

TYPE: amino acid

30 **(D)** 

TOPOLOGY: linear

(ii)

```
MOLECULE TYPE: protein
    (vi)
    ORIGINAL SOURCE
    (A)
    ORGANISM: equine infectious anemia virus
    (ix)
    FEATURE:
10
    (A)
    NAME: p26
    (x)
    PUBLICATION INFORMATION
    (A)
    AUTHORS:
15
    (B)
    TITLE: (
    C)
    JOURNAL:
    (D)
20
    VOLUME:
    (F)
    PAGES:
    (G)
    DATE:
25
    (xi)
    SEQUENCE DESCRIPTION: SEQ ID NO:1
    His His His His His Gly Ser Pro Gly Asn Pro Leu Thr Trp
                  5
                                  10
                                                    15
30
```

C 18 3

	Ser Lys Ala Leu Lys Lys Leu Glu Lys Val Thr Val Gln Gly Ser			
	20	25	30	
	Gln Lys Leu Thr Thr Gly Asn Cys Na Trp Ala Leu Ser Leu Val			
	35	40	45	
5	Asp Leu Phe His Asp Th	r Asn Phe Val Lys Glu Lys	s Asp Trp Gln	
	50	55	60	
	Leu Arg Asp Val IIe Pro I	Leu Leu Glu Asp Val Thr (	Gln Thr Val	
	65	70	75	
	Ser Gly Gln Glu Arg Glu	Ala Phe Glu Arg Thr Trp	Trp Ala Ile	
10	80	85	90	
	Ser Ala Val Lys Met Gly	Leu Gln Ile Asn AsnVal Va	al Asp Gly	
	95	100	105	
	Lys Ala Ser Phe Gln Leu Leu Arg Ala Lys Tyr Glu Lys Lys Thr			
	110	115	120	
15	Ala Asn Lys Lys Gln Ser	Glu Pro Ser Glu Glu Tyr F	Pro Ile Met	
	125	130	135	
	lle Asp Gly Ala Gly Asn A	Arg Asn Phe Arg Pro Leu	Thr Pro Arg	
	140	145	150	
	Gly Tyr Thr Thr Trp Val A	AsnThr Ile Gin Thr Asn Gly	Leu Leu	
20	155	160	165	
	Asn Glu Ala Ser Gln Asn	Leu Phe Gly Ile Leu Ser \	√al Asp Cys	
	170	175	180	
	Thr Ser Glu Glu Met Asn	Ala Phe Leu Asp Val Val	Pro Gly Gln	
	185	190	195	
25	Ala Gly Gln Lys Gln lle Le	eu Leu Asp Ala Ile Asp Ly	s Ile Ala	
	200	205	210	
	Asp Asp Trp Asp Asn Arg	j His Pro Leu Pro Asn Ala	Pro Leu Val	
	215	220	225	
	Ala Pro Pro Gin Gly Pro I	le Pro Met Thr Ala Arg Ph	e Ile Arg	
30	230	235	240	
	Gly Leu Gly Val Pro Arg (	Glu Arg Gln Met Glu Pro		
	245	250		

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	Asn Cys Val Val Gln Ser Phe Gly Val Ile Gly Gln Ala His Leu.			
	260	265	270	
	Glu Leu Pro Arg Pro Asn Lys Arg Ile Arg Asn Gln. Ser Phe Asn			
	275	280	285	
5	Gln Tyr Asn Cys Ser Ile Asn. Asn Lys Thr Glu Leu Glu Thr Trp			
	290	295	300	
	Lys Leu.Val Lys Thr Ser Gly Val Thr Pro Leu Pro. lle Ser Ser			
	305	310	315	
	Glu Ala Asn Thr Gly Leu			
10	320			